

PDB code

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graph TD; A[/PDB code/] -.-> B[Map code to AlphaFold equivalent]; A --> C[Fetch PDB files]; B --> C; C --> D[Domain assignment]; D --> E[Evaluate];
```

The flowchart illustrates a five-step process for protein structure analysis. It begins with a parallelogram box labeled 'PDB code'. A solid arrow points down from this box to a large rounded rectangle labeled 'Fetch PDB files'. A dotted arrow points right from the 'PDB code' box to another rounded rectangle labeled 'Map code to AlphaFold equivalent'. A solid arrow points down from this second box to the 'Fetch PDB files' box. From 'Fetch PDB files', a solid arrow points down to a rounded rectangle labeled 'Domain assignment'. Finally, a solid arrow points down from 'Domain assignment' to the last rounded rectangle, labeled 'Evaluate'.

Map code to
AlphaFold equivalent

Fetch PDB files

Domain assignment

Evaluate